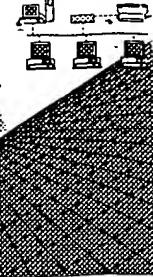


BIOTECHNOLOGY
SYSTEMS
BRANCH



16C1

1647

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/613,591B
Source: O/PE
Date Processed by STIC: 6/5/2001

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TECH CENTER 1600/2900

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

R. Deitz

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001

TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt
Output Set: C:\CRF3\06052001\I613591B.raw

3 <110> APPLICANT: BOYLE, WILLIAM
4 LACEY, DAVID
5 CALZONE, FRANK
6 CHANG, MING-SHI
7 SENALDI, GIORGIO
9 <120> TITLE OF INVENTION: COMBINATION THERAPY FOR CONDITIONS LEADING TO BONE LOSS
11 <130> FILE REFERENCE: A-378CIP5
13 <140> CURRENT APPLICATION NUMBER: US 09/613,591B
14 <141> CURRENT FILING DATE: 2000-07-10
16 <150> PRIOR APPLICATION NUMBER: US 09/457,647
17 <151> PRIOR FILING DATE: 1999-12-09
19 <150> PRIOR APPLICATION NUMBER: US 09/350,670
20 <151> PRIOR FILING DATE: 1999-07-09
22 <150> PRIOR APPLICATION NUMBER: US 08/706,945
23 <151> PRIOR FILING DATE: 1996-09-03
25 <150> PRIOR APPLICATION NUMBER: US 08/577,788
26 <151> PRIOR FILING DATE: 1995-12-22
28 <160> NUMBER OF SEQ ID NOS: 168
30 <170> SOFTWARE: PatentIn version 3.0
32 <210> SEQ ID NO: 1
33 <211> LENGTH: 36
34 <212> TYPE: DNA
35 <213> ORGANISM: Artificial
37 <220> FEATURE:
38 <223> OTHER INFORMATION: Random cDNA with internal NotI restriction site.
40 <220> FEATURE:
41 <221> NAME/KEY: misc_feature
42 <222> LOCATION: (16, 17, 18, 19, 20, 21, 22 and) .. (23) *h's are not at these locations* *(28)-(35) these are the correct locations*
43 <223> OTHER INFORMATION: N = any random nucleic acid
46 <400> SEQUENCE: 1
W--> 47 aaaggaagga aaaaagcggc cgctacannn nnnnnnt 36
50 <210> SEQ ID NO: 2
51 <211> LENGTH: 16
52 <212> TYPE: DNA
OK-> 53 <213> ORGANISM: Artificial
55 <220> FEATURE:
56 <223> OTHER INFORMATION: ds oligonucleotide adapter
58 <400> SEQUENCE: 2
59 tcgaccacg cgtccg 16
62 <210> SEQ ID NO: 3
63 <211> LENGTH: 12
64 <212> TYPE: DNA
OK-> 65 <213> ORGANISM: Artificial
67 <220> FEATURE:
68 <223> OTHER INFORMATION: ds oligonucleotide adapter
70 <400> SEQUENCE: 3
71 gggtgcgcag gc 12

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001
TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt
Output Set: C:\CRF3\06052001\I613591B.raw

74 <210> SEQ ID NO: 4
 75 <211> LENGTH: 18
 76 <212> TYPE: DNA
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 79 <220> FEATURE:
 80 <223> OTHER INFORMATION: PCR primer
 82 <400> SEQUENCE: 4
 83 tgtaaaacga cggccagt 18
 86 <210> SEQ ID NO: 5
 87 <211> LENGTH: 18
 88 <212> TYPE: DNA
~~OK~~ 89 <213> ORGANISM: Artificial
 91 <220> FEATURE:
 92 <223> OTHER INFORMATION: PCR primer
 94 <400> SEQUENCE: 5
 95 caggaaacag ctatgacc 18
 98 <210> SEQ ID NO: 6
 99 <211> LENGTH: 20
 100 <212> TYPE: DNA
~~OK~~ 101 <213> ORGANISM: Artificial
 103 <220> FEATURE:
 104 <223> OTHER INFORMATION: T3 primer
 106 <400> SEQUENCE: 6
 107 caatttaaccc tcactaaagg 20
 110 <210> SEQ ID NO: 7
 111 <211> LENGTH: 23
 112 <212> TYPE: DNA
 113 <213> ORGANISM: Rattus rattus
 115 <400> SEQUENCE: 7
 116 gcatttatgac ccagaaaaccg gac 23
 119 <210> SEQ ID NO: 8
 120 <211> LENGTH: 23
 121 <212> TYPE: DNA
 122 <213> ORGANISM: Rattus rattus
 124 <400> SEQUENCE: 8
 125 aggttagcgcc cttcctcaca ttc 23
 128 <210> SEQ ID NO: 9
 129 <211> LENGTH: 30
 130 <212> TYPE: DNA
~~OK~~ 131 <213> ORGANISM: Artificial
 133 <220> FEATURE:
 134 <223> OTHER INFORMATION: PCR primer
 136 <400> SEQUENCE: 9
 137 gactagtccc acaatgaaca agtggctgtg 30
 140 <210> SEQ ID NO: 10
 141 <211> LENGTH: 45
 142 <212> TYPE: DNA
~~OK~~ 143 <213> ORGANISM: Artificial
 145 <220> FEATURE:

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001
 TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt
 Output Set: C:\CRF3\06052001\I613591B.raw

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146 <223> OTHER INFORMATION: PCR primer
148 <400> SEQUENCE: 10
149 ataagaatgc ggccgctaaa ctatgaaaca gcccagtgtac cattc          45
152 <210> SEQ ID NO: 11
153 <211> LENGTH: 21
154 <212> TYPE: DNA
OK 155 <213> ORGANISM: Artificial
157 <220> FEATURE:
158 <223> OTHER INFORMATION: PCR primer
160 <400> SEQUENCE: 11
161 gcctctagaa agagctggga c                                         21
164 <210> SEQ ID NO: 12
165 <211> LENGTH: 21
166 <212> TYPE: DNA
OK 167 <213> ORGANISM: Artificial
169 <220> FEATURE:
170 <223> OTHER INFORMATION: PCR primer
172 <400> SEQUENCE: 12
173 cgccgtgttc catttatgag c                                         21
176 <210> SEQ ID NO: 13
177 <211> LENGTH: 24
178 <212> TYPE: DNA
179 <213> ORGANISM: Rattus rattus
181 <400> SEQUENCE: 13
182 atcaaaggca gggcatactt cctg                                     24
185 <210> SEQ ID NO: 14
186 <211> LENGTH: 24
187 <212> TYPE: DNA
188 <213> ORGANISM: Rattus rattus
190 <400> SEQUENCE: 14
191 gttgcactcc tgtttcacgg tctg                                     24
194 <210> SEQ ID NO: 15
195 <211> LENGTH: 24
196 <212> TYPE: DNA
197 <213> ORGANISM: Rattus rattus
199 <400> SEQUENCE: 15
200 caagacaccc tgaaggccct gatg                                     24
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204 <211> LENGTH: 24
205 <212> TYPE: DNA
206 <213> ORGANISM: Rattus rattus
208 <400> SEQUENCE: 16
209 taactttac agaagagcat cagc                                     24
212 <210> SEQ ID NO: 17
213 <211> LENGTH: 33
214 <212> TYPE: DNA
215 <213> ORGANISM: Rattus rattus
217 <400> SEQUENCE: 17
218 agcgccggccg catgaacaag tggctgtgct gcg                         33

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001
TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt
Output Set: C:\CRF3\06052001\I613591B.raw

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221 <210> SEQ ID NO: 18
222 <211> LENGTH: 31
223 <212> TYPE: DNA
224 <213> ORGANISM: Rattus rattus
226 <400> SEQUENCE: 18
227 agctctagag aaacagccca gtgaccattc c                                31
230 <210> SEQ ID NO: 19
231 <211> LENGTH: 24
232 <212> TYPE: DNA
233 <213> ORGANISM: Rattus rattus
235 <400> SEQUENCE: 19
236 gtgaagctgt gcaagaacct gatg                                         24
239 <210> SEQ ID NO: 20
240 <211> LENGTH: 24
241 <212> TYPE: DNA
242 <213> ORGANISM: Rattus rattus
244 <400> SEQUENCE: 20
245 atcaaaggca gggcatactt cctg                                         24
248 <210> SEQ ID NO: 21
249 <211> LENGTH: 24
250 <212> TYPE: DNA
251 <213> ORGANISM: Homo sapiens
253 <400> SEQUENCE: 21
254 cagatcctga agctgcttag tttg                                         24
257 <210> SEQ ID NO: 22
258 <211> LENGTH: 33
259 <212> TYPE: DNA
260 <213> ORGANISM: Homo sapiens
262 <400> SEQUENCE: 22
263 agcgccgcggcccg cggggaccac aatgaacaag ttg                           33
266 <210> SEQ ID NO: 23
267 <211> LENGTH: 33
268 <212> TYPE: DNA
269 <213> ORGANISM: Homo sapiens
271 <400> SEQUENCE: 23
272 agctctagaa ttgtgaggaa acagctcaat ggc                               33
275 <210> SEQ ID NO: 24
276 <211> LENGTH: 39
277 <212> TYPE: DNA
278 <213> ORGANISM: Artificial
280 <220> FEATURE:
281 <223> OTHER INFORMATION: PCR primer
283 <400> SEQUENCE: 24
284 atagcgccgcggcccg ctgagccaa atcttgtgac aaaactcac                     39
287 <210> SEQ ID NO: 25
288 <211> LENGTH: 45
289 <212> TYPE: DNA
290 <213> ORGANISM: Artificial
292 <220> FEATURE:

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001

TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt
 Output Set: C:\CRF3\06052001\I613591B.raw

293 <223> OTHER INFORMATION: PCR primer
 295 <400> SEQUENCE: 25
 296 tcttagagtctg acttattcatt tacccggaga cagggagagg ctctt 45
 299 <210> SEQ ID NO: 26
 300 <211> LENGTH: 38
 301 <212> TYPE: DNA
 302 <213> ORGANISM: Mus musculus
 304 <400> SEQUENCE: 26
 305 cctctgagct caagcttccg aggaccacaa tgaacaag 38
 308 <210> SEQ ID NO: 27
 309 <211> LENGTH: 43
 310 <212> TYPE: DNA
 311 <213> ORGANISM: Mus musculus
 313 <400> SEQUENCE: 27
 314 cctctgcgac cgctaaggcag cttatcccac ccgttggac ctg 43
 317 <210> SEQ ID NO: 28
 318 <211> LENGTH: 38
 319 <212> TYPE: DNA
 320 <213> ORGANISM: Mus musculus
 322 <400> SEQUENCE: 28
 323 cctctgagct caagcttccg aggaccacaa tgaacaag 38
 326 <210> SEQ ID NO: 29
 327 <211> LENGTH: 24
 328 <212> TYPE: DNA
 329 <213> ORGANISM: Homo sapiens
 331 <400> SEQUENCE: 29
 332 tccgttaagaa acagccccgt gacc 24
 335 <210> SEQ ID NO: 30
 336 <211> LENGTH: 31
 337 <212> TYPE: DNA
 338 <213> ORGANISM: Mus musculus
 340 <400> SEQUENCE: 30
 341 cctctgcggc cgctgttgca tttcctttct g 31
 344 <210> SEQ ID NO: 31
 345 <211> LENGTH: 19
 346 <212> TYPE: PRT
 347 <213> ORGANISM: Mus musculus
 349 <400> SEQUENCE: 31
 351 Glu Thr Leu Pro Pro Lys Tyr Leu His Tyr Asp Pro Glu Thr Gly His
 352 1 5 10 15
 354 Gln Leu Leu
 357 <210> SEQ ID NO: 32
 358 <211> LENGTH: 21
 359 <212> TYPE: DNA
 360 <213> ORGANISM: Mus musculus
 362 <400> SEQUENCE: 32
 363 tcccttgccc tgaccactct t 21
 366 <210> SEQ ID NO: 33
 367 <211> LENGTH: 34

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001

TIME: 12:57:13

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001

TIME: 12:57:13

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

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